## AMENDMENT TO THE SPECIFICATION

Please delete Tables 1 and 2 in their entirety.

Please insert replacement Tables 1 and 2, and the following paragraphs at the end of the material added at pages 6-13 of the response filed August 24, 2009.

Table 1

| GENE | PRECURSOR-SEQUENCE   | P<br>-SEQ | GENE-SEQ   | G<br>-SEQID |              | PRECURSOR  |                  |      |
|------|----------------------|-----------|------------|-------------|--------------|------------|------------------|------|
|      |                      |           |            |             |              |            |                  |      |
| GAM  | CCCGTGGGGGGTCTTAGTGG | 1917      | TAGCACCGCT | 4642        | T G T        | CT -       | AC A             | A    |
| 1931 | AAGTGACGTGCTGTGAATAC |           | ATCCACTATG |             | CCCG GG GGG  | TAGTGGA A  | GTG GTGCTGTG A   | AT C |
|      | AGGTCCATAGCACCGCTATC |           | TCTC       |             | 1111 11 1111 | 1111111111 | 111 111111111111 | 1    |
|      | CACTATGTCTCGCCCGGG   |           |            |             | GGGC CC CTC  | ATCACCT TO | CGC CACGATAC T   | G A  |
|      |                      |           |            |             | - G T        | 'GT A      | C                | G    |

Nucleotide sequence of the VGAM PRECURSOR RNA, and of the 'diced' VGAM RNA, and a Schematic representation of the secondary folding of VGAM FOLDED PRECURSOR RNA of each of the plurality VGAM GENEs described by Fig.1 are further described hereinbelow with reference to Table 1.

Nucleotide sequences of the VGAM1931 precursor RNA, herein designated VGAM PRECURSOR RNA, and of the diced VGAM1931 RNA, herein designated VGAM RNA, and a schematic representation of the secondary folding of VGAM1931 folded precursor RNA, herein designated VGAM FOLDED PRECURSOR RNA, of VGAM1931 are further described hereinbelow with refernce to Table 1.

Table 2

| GE | NE | TARGET | UTR | SEQUENCE               | SEQID | BINDING-SITE          |
|----|----|--------|-----|------------------------|-------|-----------------------|
| == |    |        | === |                        |       |                       |
| GA | M  | COL6A1 | 3 * | AGACCCTCGAGATTAACGGTGC | 7584  | CT_ CACTAT            |
| 19 | 31 |        |     | TA                     |       | TAGCACCG ATC GTCT     |
|    |    |        |     |                        |       | 11111111 111 1111     |
|    |    |        |     |                        |       | ATCGTGGC TAG CAGA     |
|    |    |        |     |                        |       | AAT AGCTCC            |
| GA | M  | HIP12  | 3 * | GAGACATAGCGGCCCGGGCGCT | 32922 | A_ CTAT A             |
| 19 | 31 |        |     | G                      |       | TAGC CCG CC CTATGTCTC |
|    |    |        |     |                        |       |                       |
|    |    |        |     |                        |       | GTCG GGC GG GATACAGAG |
|    |    |        |     |                        |       | CG CC C               |

| GAM SFRS1 3               | • GACTTAGGTGGGTAGCAATGC      | 13801          | CC AT_ GCA GCTATCCACT GTC                   CGT CGATGGGTGG CAG AA ATT              |
|---------------------------|------------------------------|----------------|--|
| GAM FLJ2 3'<br>1931 0436  | AGACACAGTGGATATTTTGGGC       | 19472          | A C A GC CGG TATCCACT TGTCT  |
|                           | GAGACATAATGACAGCTGATGT<br>TA |                | C _ A CAC TAGGA C GCT TC TATGTCTC  |
| GAM ZNF 3'                | GAGAGCAGGGGACACGGTGCTA 1     |                | CTA ACTA _ TAGGACCG TCC TC TCTC  |
| GAM LOC14 5'<br>1931 5761 | GAGACACAGTGAAAAGGGGT 4       |                | G ATC A ACC CT CACT TETCTC                          TEG GA GTGA ACAGAG G AAA C     |
| GAM LOC14 5'<br>1931 6603 | ACATAGTGGACAGCATTGT          | GC<br>  <br>TG | CC A A GCT TCCACTATGT  |
| GAM LOC20 3'              | GAGACATAACGATTTGGATTGC<br>TA | T              | AGCA CCG ATC TATGTCTC  |
| GAM LOC5 5'               | GAAATATGGATAGCAGC            |                | ACC C G GC GCTATCCA TAT TC   |
| GAM 1005 3'               | GAGATATAGCAACTAACGACGC<br>TA |                | AC C TCCA TAGC CG TA CTATGTCTC                      ATCG GC AT GATATAGAG CA A CAAC |

Nucleotide sequence of host target binding sites, such as BINDING SITE-I,BINDING SITE-II and BINDING SITE-III of Fig.1, found on, and schematic representation of the complementarity of each of these host target binding sites to VGAM RNA are described hereinbelow with reference to Table 2.

Nucleotide sequences of host target binding sites, such as BINDING SITE-I, BINDING SITE-III and BINDING SITE-III of Fig.1, found on, and schematic representation of the

complementarity of each of these host target binding sites to VGAM1931 RNA, herein designated VGAM RNA, are described hereinbelow with reference to Table 2.